



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/660,118
Source: oipe
Date Processed by STIC: 9-24-03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to:
U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
Or
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 04/24/2003

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER: 10/660,118

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
- 2 Invalid Line Length The rules require that a line **not exceed** 72 characters in length. This includes white spaces.
- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do **not** use tab codes between numbers; use **space characters**, instead.
- 4 Non-ASCII The submitted file was **not** saved in ASCII(DOS) text, as **required** by the Sequence Rules. Please **ensure your subsequent submission is saved in ASCII text**.
- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. **Per Sequence Rules, each n or Xaa can only represent a single residue.** Please present the **maximum** number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. **This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.**
- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for **each** skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000
- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is **MANDATORY** if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only **valid** <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is **required** when <213> response is Unknown or is Artificial Sequence
- 11 X Use of <220> Sequence(s) 1-3 missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is **MANDATORY** if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/660,118

DATE: 09/24/2003

TIME: 10:37:18

Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

3 <110> APPLICANT: White, Carl W.
 5 <120> TITLE OF INVENTION: Product and Process for Liquefaction of Mucus or Sputum
 7 <130> FILE REFERENCE: 2879-98
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/10/660,118
 C--> 9 <141> CURRENT FILING DATE: 2003-09-10
 9 <150> PRIOR APPLICATION NUMBER: 60/409,960
 10 <151> PRIOR FILING DATE: 2002-09-10
 12 <150> PRIOR APPLICATION NUMBER: 60/462,082
 13 <151> PRIOR FILING DATE: 2003-04-11
 15 <160> NUMBER OF SEQ ID NOS: 15
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 4
 21 <212> TYPE: PRT
 22 <213> ORGANISM: Artificial sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: peptide insufficient explanation
 27 <400> SEQUENCE: 1 give source of genetic material
 29 Cys Gly Pro Cys see item 11 on error summary
 30 1 report.
 33 <210> SEQ ID NO: 2
 34 <211> LENGTH: 6
 35 <212> TYPE: PRT
 36 <213> ORGANISM: Artificial sequence
 38 <220> FEATURE:
 39 <223> OTHER INFORMATION: peptide
 41 <220> FEATURE:
 42 <221> NAME/KEY: misc_feature
 43 <222> LOCATION: (1)..(6)
 44 <223> OTHER INFORMATION: Xaa = any amino acid
 47 <400> SEQUENCE: 2
 W--> 49 Xaa Cys Gly Pro Cys Xaa
 50 1 5
 53 <210> SEQ ID NO: 3
 54 <211> LENGTH: 6
 55 <212> TYPE: PRT
 56 <213> ORGANISM: Artificial sequence
 58 <220> FEATURE:
 59 <223> OTHER INFORMATION: peptide
 61 <400> SEQUENCE: 3
 63 Trp Cys Gly Pro Cys Lys
 64 1 5
 67 <210> SEQ ID NO: 4

RAW SEQUENCE LISTING

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TIME: 10:37:18

Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

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68 <211> LENGTH: 109
69 <212> TYPE: PRT
70 <213> ORGANISM: Pseudomonas syringae
72 <400> SEQUENCE: 4
74 Met Ser Asn Asp Leu Ile Lys His Val Thr Asp Ala Ser Phe Glu Ala
75 1 5 10 15
78 Asp Val Leu Lys Ala Asp Gly Ala Val Leu Val Asp Tyr Trp Ala Glu
79 20 25 30
82 Trp Cys Gly Pro Cys Lys Met Ile Ala Pro Val Leu Asp Glu Ile Ala
83 35 40 45
86 Thr Thr Tyr Ala Gly Lys Leu Thr Ile Ala Lys Leu Asn Ile Asp Glu
87 50 55 60
90 Asn Gln Glu Thr Pro Ala Lys His Gly Val Arg Gly Ile Pro Thr Leu
91 65 70 75 80
94 Met Leu Phe Lys Asn Gly Asn Val Glu Ala Thr Lys Val Gly Ala Leu
95 85 90 95
98 Ser Lys Ser Gln Leu Ala Ala Phe Leu Asp Ala Asn Ile
99 100 105
102 <210> SEQ ID NO: 5
103 <211> LENGTH: 104
104 <212> TYPE: PRT
105 <213> ORGANISM: Porphyromonas gingivalis
107 <400> SEQUENCE: 5
109 Met Ala Leu Gln Ile Thr Asp Ala Thr Phe Asp Gly Leu Val Ala Glu
110 1 5 10 15
113 Gly Lys Pro Met Val Val Asp Phe Trp Ala Thr Trp Cys Gly Pro Cys
114 20 25 30
117 Arg Met Val Gly Pro Ile Ile Asp Glu Leu Ala Ala Glu Tyr Glu Gly
118 35 40 45
121 Arg Ala Ile Ile Gly Lys Val Asp Val Asp Ala Asn Thr Glu Leu Pro
122 50 55 60
125 Met Lys Tyr Gly Val Arg Asn Ile Pro Thr Ile Leu Phe Ile Lys Asn
126 65 70 75 80
129 Gly Glu Val Val Lys Lys Leu Val Gly Ala Gln Ser Lys Asp Val Phe
130 85 90 95
133 Lys Lys Glu Leu Asp Ala Leu Phe
134 100
137 <210> SEQ ID NO: 6
138 <211> LENGTH: 103
139 <212> TYPE: PRT
140 <213> ORGANISM: Listeria monocytogenes
142 <400> SEQUENCE: 6
144 Met Val Lys Glu Ile Thr Asp Ala Thr Phe Glu Gln Glu Thr Ser Glu
145 1 5 10 15
148 Gly Leu Val Leu Thr Asp Phe Trp Ala Thr Trp Cys Gly Pro Cys Arg
149 20 25 30
152 Met Val Ala Pro Val Leu Glu Glu Ile Gln Glu Glu Arg Gly Glu Ala
153 35 40 45
156 Leu Lys Ile Val Lys Met Asp Val Asp Glu Asn Pro Glu Thr Pro Gly

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Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

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157      50      55      60
160 Ser Phe Gly Val Met Ser Ile Pro Thr Leu Leu Ile Lys Lys Asp Gly
161 65      70      75      80
164 Glu Val Val Glu Thr Ile Ile Gly Tyr Arg Pro Lys Glu Glu Leu Asp
165      85      90      95
168 Glu Val Ile Asn Lys Tyr Val
169      100
172 <210> SEQ ID NO: 7
173 <211> LENGTH: 103
174 <212> TYPE: PRT
175 <213> ORGANISM: Saccharomyces cerevisiae
177 <400> SEQUENCE: 7
179 Met Val Thr Gln Phe Lys Thr Ala Ser Glu Phe Asp Ser Ala Ile Ala
180 1      5      10      15
183 Gln Asp Lys Leu Val Val Val Asp Phe Tyr Ala Thr Trp Cys Gly Pro
184      20      25      30
187 Cys Lys Met Ile Ala Pro Met Ile Glu Lys Phe Ser Glu Gln Tyr Pro
188      35      40      45
191 Gln Ala Asp Phe Tyr Lys Leu Asp Val Asp Glu Leu Gly Asp Val Ala
192      50      55      60
195 Gln Lys Asn Glu Val Ser Ala Met Pro Thr Leu Leu Leu Phe Lys Asn
196 65      70      75      80
199 Gly Lys Glu Val Ala Lys Val Val Gly Ala Asn Pro Ala Ala Ile Lys
200      85      90      95
203 Gln Ala Ile Ala Ala Asn Ala
204      100
207 <210> SEQ ID NO: 8
208 <211> LENGTH: 105
209 <212> TYPE: PRT
210 <213> ORGANISM: Gallus gallus
212 <400> SEQUENCE: 8
214 Met Val Lys Ser Val Gly Asn Leu Ala Asp Phe Glu Ala Glu Leu Lys
215 1      5      10      15
218 Ala Ala Gly Glu Lys Leu Val Val Val Asp Phe Ser Ala Thr Trp Cys
219      20      25      30
222 Gly Pro Cys Lys Met Ile Lys Pro Phe Phe His Ser Leu Cys Asp Lys
223      35      40      45
226 Phe Gly Asp Val Val Phe Ile Glu Ile Asp Val Asp Asp Ala Gln Asp
227      50      55      60
230 Val Ala Thr His Cys Asp Val Lys Cys Met Pro Thr Phe Gln Phe Tyr
231 65      70      75      80
234 Lys Asn Gly Lys Lys Val Gln Glu Phe Ser Gly Ala Asn Lys Glu Lys
235      85      90      95
238 Leu Glu Glu Thr Ile Lys Ser Leu Val
239      100      105
242 <210> SEQ ID NO: 9
243 <211> LENGTH: 105
244 <212> TYPE: PRT
245 <213> ORGANISM: Mus musculus

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Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

247 <400> SEQUENCE: 9

249 Met Val Lys Leu Ile Glu Ser Lys Glu Ala Phe Gln Glu Ala Leu Ala
 250 1 5 10 15
 253 Ala Ala Gly Asp Lys Leu Val Val Val Asp Phe Ser Ala Thr Trp Cys
 254 20 25 30
 257 Gly Pro Cys Lys Met Ile Lys Pro Phe Phe His Ser Leu Cys Asp Lys
 258 35 40 45
 261 Tyr Ser Asn Val Val Phe Leu Glu Val Asp Val Asp Cys Gln Asp
 262 50 55 60
 265 Val Ala Ala Asp Cys Glu Val Lys Cys Met Pro Thr Phe Gln Phe Tyr
 266 65 70 75 80
 269 Lys Lys Gly Gln Lys Val Gly Glu Phe Ser Gly Ala Asn Lys Glu Lys
 270 85 90 95
 273 Leu Glu Ala Ser Ile Thr Glu Tyr Ala
 274 100 105

277 <210> SEQ ID NO: 10

278 <211> LENGTH: 105

279 <212> TYPE: PRT

280 <213> ORGANISM: Rattus norvegicus

282 <400> SEQUENCE: 10

284 Met Val Lys Leu Ile Glu Ser Lys Glu Ala Phe Gln Glu Ala Leu Ala
 285 1 5 10 15
 288 Ala Ala Gly Asp Lys Leu Val Val Val Asp Phe Ser Ala Thr Trp Cys
 289 20 25 30
 292 Gly Pro Cys Lys Met Ile Lys Pro Phe Phe His Ser Leu Cys Asp Lys
 293 35 40 45
 296 Tyr Ser Asn Val Val Phe Leu Glu Val Asp Val Asp Asp Cys Gln Asp
 297 50 55 60
 300 Val Ala Ala Asp Cys Glu Val Lys Cys Met Pro Thr Phe Gln Phe Tyr
 301 65 70 75 80
 304 Lys Lys Gly Gln Lys Val Gly Glu Phe Ser Gly Ala Asn Lys Glu Lys
 305 85 90 95
 308 Leu Glu Ala Thr Ile Thr Glu Phe Ala
 309 100 105

312 <210> SEQ ID NO: 11

313 <211> LENGTH: 105

314 <212> TYPE: PRT

315 <213> ORGANISM: Bos taurus

317 <400> SEQUENCE: 11

319 Met Val Lys Gln Ile Glu Ser Lys Tyr Ala Phe Gln Glu Ala Leu Asn
 320 1 5 10 15
 323 Ser Ala Gly Glu Lys Leu Val Val Val Asp Phe Ser Ala Thr Trp Cys
 324 20 25 30
 327 Gly Pro Cys Lys Met Ile Lys Pro Phe Phe His Ser Leu Ser Glu Lys
 328 35 40 45
 331 Tyr Ser Asn Val Val Phe Leu Glu Val Asp Val Asp Asp Cys Gln Asp
 332 50 55 60
 335 Val Ala Ala Glu Cys Glu Val Lys Cys Met Pro Thr Phe Gln Phe Phe
 336 65 70 75 80

RAW SEQUENCE LISTING

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DATE: 09/24/2003

TIME: 10:37:18

Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

```

339 Lys Lys Gly Gln Lys Val Gly Glu Phe Ser Gly Ala Asn Lys Glu Lys
340      85      90      95
343 Leu Glu Ala Thr Ile Asn Glu Leu Ile
344      100      105
347 <210> SEQ ID NO: 12
348 <211> LENGTH: 105
349 <212> TYPE: PRT
350 <213> ORGANISM: Homo sapiens
352 <400> SEQUENCE: 12
354 Met Val Lys Gln Ile Glu Ser Lys Thr Ala Phe Gln Glu Ala Leu Asp
355 1      5      10      15
358 Ala Ala Gly Asp Lys Leu Val Val Val Asp Phe Ser Ala Thr Trp Cys
359      20      25      30
362 Gly Pro Cys Lys Met Ile Lys Pro Phe Phe His Ser Leu Ser Glu Lys
363      35      40      45
366 Tyr Ser Asn Val Ile Phe Leu Glu Val Asp Val Asp Asp Cys Gln Asp
367      50      55      60
370 Val Ala Ser Glu Cys Glu Val Lys Cys Met Pro Thr Phe Gln Phe Phe
371 65      70      75      80
374 Lys Lys Gly Gln Lys Val Gly Glu Phe Ser Gly Ala Asn Lys Glu Lys
375      85      90      95
378 Leu Glu Ala Thr Ile Asn Glu Leu Val
379      100      105
382 <210> SEQ ID NO: 13
383 <211> LENGTH: 134
384 <212> TYPE: PRT
385 <213> ORGANISM: Arabidopsis thaliana
387 <400> SEQUENCE: 13
389 Met Gly Gly Ala Leu Ser Thr Val Phe Gly Ser Gly Glu Asp Ala Ala
390 1      5      10      15
393 Ala Ala Gly Thr Glu Ser Ser Glu Pro Ser Arg Val Leu Lys Phe Ser
394      20      25      30
397 Ser Ser Ala Arg Trp Gln Leu His Phe Asn Glu Ile Lys Glu Ser Asn
398      35      40      45
401 Lys Leu Leu Val Val Asp Phe Ser Ala Ser Trp Cys Gly Pro Cys Arg
402      50      55      60
405 Met Ile Glu Pro Ala Ile His Ala Met Ala Asp Lys Phe Asn Asp Val
406 65      70      75      80
409 Asp Phe Val Lys Leu Asp Val Asp Glu Leu Pro Asp Val Ala Lys Glu
410      85      90      95
413 Phe Asn Val Thr Ala Met Pro Thr Phe Val Leu Val Lys Arg Gly Lys
414      100      105      110
417 Glu Ile Glu Arg Ile Ile Gly Ala Lys Lys Asp Glu Leu Glu Lys Lys
418      115      120      125
421 Val Ser Lys Leu Arg Ala
422      130
425 <210> SEQ ID NO: 14
426 <211> LENGTH: 167
427 <212> TYPE: PRT

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/660,118

DATE: 09/24/2003
TIME: 10:37:19

Input Set : A:\2879-98.ST25.txt
Output Set: N:\CRF4\09242003\J660118.raw

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; Xaa Pos. 1,6

VERIFICATION SUMMARY

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DATE: 09/24/2003

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Input Set : A:\2879-98.ST25.txt

Output Set: N:\CRF4\09242003\J660118.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application No
L:9 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:49 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:0